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Glenn P. Ladwig, Patent Attorney

SUPPLEMENTAL PRELIMINARY

AMENDMENT

Examining Group 1645

Patent Application

Docket No. GJE-65

Serial No. 09/830,807

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Examiner

Jana A. Hines

Art Unit

1645

Applicants

Helen Rachel Crooke, Enda Elizabeth Clarke, Paul Howard Everest, Gordon Dougan, David William Holden, Jacqueline Elizabeth Shea,

Robert Graham Feldman

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2112

For

Virulence Genes and Proteins, and Their Use

Assistant Commissioner for Patents

Washington, D.C. 20231

SUPPLEMENTAL PRELIMINARY AMENDMENT

Sir:

Applicants respectfully request that the above-identified patent application be amended as follows:

In the Claims

Please substitute the following claims:

Claim 1 (twice amended):

1. An isolated peptide encoded by an operon, wherein said operon comprises a gene selected from the group consisting of tatA, tatB, tatC, tatD, tatE, mdoG, creC, recG, yggN, eck1, iroD, iroC, iroE, mtd2, and ms1 to ms16, obtainable from E. coli K1, or a homologue thereof in a Gram-negative bacterium, wherein said homologue has at least 30% homology at the amino acid or nucleotide level, or a functional fragment thereof.



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Claim 3 (twice amended):

3. An isolated polynucleotide which comprises a gene selected from the group consisting of tatA, tatB, tatC, tatD, tatE, mdoG, creC, recG, yggN, eck1, iroD, iroC, iroE, mtd2, and ms1 to ms16, obtainable from E. coli K1, or a homologue thereof in a Gram-negative bacterium, wherein said homologue has at least 30% homology at the amino acid or nucleotide level, or a functional fragment thereof.

Claim 4 (twice amended):

4. A host transformed to express a peptide encoded by an operon, wherein said operon comprises a gene selected from the group consisting of tatA, tatB, tatC, tatD, tatE, mdoG, creC, recG, yggN, eck1, iroD, iroC, iroE, mtd2, and ms1 to ms16, obtainable from E. coli K1, or a homologue thereof in a Gram-negative bacterium, wherein said homologue has at least 30% homology at the amino acid or nucleotide level, or a functional fragment thereof.

Claim 5 (twice amended):

5. A vaccine comprising a peptide, or the means for its expression, wherein said peptide is encoded by an operon, wherein said operon comprises a gene selected from the group consisting of tatA, tatB, tatC, tatD, tatE, mdoG, creC, recG, yggN, eck1. iroD, iroC, iroE, mtd2, and ms1 to ms16, obtainable from E. coli K1, or a homologue thereof in a Gram-negative bacterium, wherein said homologue has at least 30% homology at the amino acid or nucleotide level, or a functional fragment thereof.

Claim 6 (twice amended):

6. A vaccine comprising a microorganism having a virulence gene mutation, wherein the gene is selected from the group consisting of tatA, tatB, tatC, tatD, tatE, mdoG, creC, recG, yggN, eck1, iroD, iroC, iroE, mtd2, and ms1 to ms16, obtainable from E. coli K1, or a homologue thereof in a Gram-negative bacterium, wherein said homologue has at least 30% homology at the amino acid or nucleotide level, or a functional fragment thereof.

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Claim 9 (twice amended):

9. A method for screening potential drugs, or for the detection of virulence, wherein said method utilizes a peptide encoded by an operon, wherein said operon comprises a gene selected from the group consisting of tatA, tatB, tatC, tatD, tatE, mdoG, creC, recG, yggN, eckl, iroD, iroC, iroE, mtd2, and ms1 to ms16, obtainable from E. coli K1, or a homologue thereof in a Gram-negative bacterium, wherein said homologue has at least 30% homology at the amino acid or nucleotide level, or a functional fragment thereof.

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Claim 10 (twice amended):

10. A method for treatment or prevention of a condition associated with infection by a Gramnegative bacterium, said method comprising administering a vaccine to a person or animal in need thereof, wherein said vaccine comprises a peptide, or a host transformed to express said peptide, wherein said peptide is encoded by an operon comprising a gene selected from the group consisting of tatA, tatB, tatD, tatC, tatE, mdoG, creC, recG, yggN, eck1, iroD, iroC, iroE, mtd2, and ms1 to ms16, obtainable from E. coli K1, or a homologue thereof in a Gram-negative bacterium, wherein said homologue has at least 30% homology at the amino acid or nucleotide level, or a functional fragment thereof.

Claim 22 (amended):

22. A method for treatment or prevention of a condition associated with infection by a Gramnegative bacterium, said method comprising administering a nucleotide to a person or animal in need thereof, wherein said nucleotide comprises an operon including a gene selected from the group consisting of tatA, tatB, tatC, tatD, tatE, mdoG, creC, recG, yggN, eck1, iroD, iroC, iroE, mtd2, and ms1 to ms16, obtainable from E. coli K1, or a homologue thereof in a Gram-negative bacterium, wherein said homologue has at least 30% homology at the amino acid or nucleotide level, or a functional fragment thereof.

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